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10 20 30 40 50 60
AACGGATCTG CCCGCCTCAG CCTCCCAAAG TGCTGGGATT GCAGGCGTGA GCCACCTCAC

70 80 90 100 110 120
CTGGCTACAA GTTTTCAAAA TACATTTATC TAGTACCCAT ACATTCTCCA GTTTGTCCAC

130 140 150 160 170 180
AGGACATCTT ATGACTTGAG CAAGCTGCTA AAAATCCAAG GGTGCAGCGT TTGTATGTCT

190 200 210 220 230 240
ATAGGATTGC TCAGATCTGC CCCCACCTG AAAGAATTTA AGAGAATTTT TTGAGGCCAG

250 260 270 280 290 300
GCACAGTGGC TCACACCTGT AATTCCAGTA CTGTGAGAGT CCGAGGTCAG AGGACTGCTT
PPRE

310 320 330 340 350 360
GAGGCCAGGA GTTCAAGAGC AGCCTGGACA ACACAGGGAG ACCTGTCACT ACAAAGAATA

370 380 390 400 410 420
AATAAATTAG CCAGGCTTAG TGGCTCATCC CTGTGGTCCC AGCTACTAGG GAGGCAGAAG

430 440 450 460 470 480
TAGGACTGCT TGTCCCAGGA GGTCAGACT GCAGTGAGCT GAGACCCAGC CACCTGCATT

490 500 510 520 530 540
CCAGCCTGGG CAACAAAAAG AGACCCTGTC TCAAAAAATA AGTTAAATAA ATAAATAATA

550 560 570 580 590 600
AAAATAGTTT AAACCCTAAA CACATCTTCT TTTTCAAAGA GGA CTCTTA AGGACTTCAT

610 620 630 640 650 660
GCTGCGTCCT GTTGATCTCC ACTTCCCTTT TTCAGCGTCC AACTTTTAA CAGTCTCTTT

FIG. 1A



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670 680 690 700 710 720
TGCCAAGGAT AATAAGTATA TAGTTTCTGG AATCCAGATT CTTCCCTGTT TGGACAGCCA

730 740 750 760 770 780
GGGGGACAAT TTTTGGTCTG CAGGCCTTTG CATCTGTTCT GCTGTTGCTC AGCAATCTCA
GRE

790 800 810 820 830 840
CAGCAAATTT GCCGAGCCTC TCCGGAATGC ACAGCCAGAC AGAGCTCAGC GCAAAAAGCTA

850 860 870 880 890 900
GAGAACCTGG CGGAGGGAGA CTCACAGTGC CACAAAAAAA CTTTATCTTT TCTTTTTTTT

910 920 930 940 950 960
TTTCTTTTCT TTCTTTCTCT TTCTTTCTTG TCTTTCTGTC TTCTCTCTCT CTCTCTCTGT

970 980 990 1000 1010 1020
CTTTCTTTCC TCTCTTTCTT TCTTTTTTCC TACATGGCAA GATCTCCTCA TGGCAGAAAT

1030 1040 1050 1060 1070 1080
AATCTGCCTT GACTTCTGTT TCCACGCTGC TTCTGCCAGG ACCATGCGCT CGGCGTGTTT
GRE

1090 1100 1110 1120 1130 1140
TTCTTTCCGC TATAATTATC CAGGCCCATC CCAGCTCTGG TCCCCTCAGC TGTTCCCTGG

1150 1160 1170 1180 1190 1200
CAGTCCCTTC TGCTGGTGAA AACACATATG GCGCCGGCCT GACCAGGGTG TAAGTGTGTG

1210 1220 1230 1240 1250 1260
AATATCAGGA AGATGACTGA ACGTCTTTGG GACTCCGTTT CCTCATTGTA AAATGGAGGT

FIG. 1B



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1270 1280 1290 1300 1310 1320
TAATACCAGC CTTCTTCTAC TCCCCAAACG CACGTGTTTG TCCCGGCCAG AGGGCCCAAT
C/EBP

1330 1340 1350 1360 1370 1380
TGTTGGCTGT TCACGCATCA GTTACCCCCA CAGGACGGGT CAGCCAATTA AAGGCGAACC
C/EBP

1390 1400 1410 1420 1430 1440
AGGCCCGGTC CATCTCCTGA CGCCTTTTCT CATCCCAGGG CTGGACAGGC AGCTGGCCTG
MyoD

1450 1460 1470 1480 1490 1500
GGCCCCGGCTC TGCCTTGTCA CGTGCGGGGG CCGGCCCCGT TGCTTGTCTG TGTGTAGGAG
GRE

1510 1520 1530 1540 1550 1560
CGTGAGGTCA CGCTGGGTGC TCCCGCCCCG CCGGGGCCTT TAGTGTCCCT GGTCCCTAAA

1570 1580 1590 1600 1610 1620
CGCCAGGCCG CTCCACCGGG GGAGAAGGCG CGAACCCAG CCGAGCCCAA CGGCTGTTGT

1630 1640 1650 1660 1670 1680
CGGTTGCCGG GCCACCTGTT GCTGCAGTTC TGATTGGTTC CTTCCCCCGA CAACGCGGCG

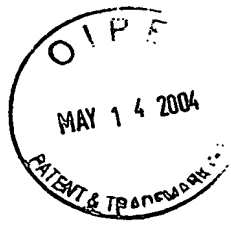
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GCTGTAACCA ATCGACAGCG AGGCCGGTCG CGAGGCCCCA GTCCCGCCCT GCAGGAGCCA
C/EBP

1750 1760 1770 1780 1790 1800
GCCGCGCGCT CGCTCGCAGG AGGGTGGGTA GTTTGCCAG CGTAGGGGGG CTGGGCCCCAT

1810 1820 1830 1840 1850 1860
AAAAGAGGAA GTGCACTTAA GACACGGCCC CGCTGGACGC TTGTTAGAAA CCGTCCTGGC

1870 1880 1890 1900 1910 1920
TGGGAAGGCA AGAGGTGTGT GACTGGACAA GACTTGTTTC TGGCGGTCAG TCTTGCCATC

FIG. 1C



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1930	1940	1950	1960	1970	1980
CTCACAGAGG TTGGCGGCCC GAGAGAGTGT GAGGCAGAGG CGGGGAGTGG CAAGGGAGTG					
1990	2000	2010	2020	2030	2040
ACCATCTCGG GGAACGAAGG AGTAAACGCG GTGATGGGAC GCACGGAAAC GGGAGTGGAG					
2050	2060	2070	2080	2090	2100
AAAGTCATGG AGAGAACCCCT AGGCGGGGCG GTCCCCGCGG AAAGGCGGCT GCTCCAGGGT					
2110	2120	2130	2140	2150	2160
CTCCGCACCC AAGTAGGAGC TGGCAGGCCC GGCCCCGCCG CGCAGGCCCC ACCCCGGGCC					
2170	2180	2190	2200	2210	2220
CCGCCCCCGA GGCTTAAGCC GCGCCGCCGC CTGCGCGGAG CCCCCTGCG AAGCCCAGCT					
2230	2240	2250	2260	2270	2280
GCGCGCGCCT TGGGATTGAC TGTCCACGCT CGCCCGGCTC GTCCGACGCG CCCTCCGCCA					
2290	2300	2310	2320	2330	2340
GCCGACAGAC ACAGCCGCAC GCACTGCCGT GTTCTCCCTG CGGCTCGGTG AGCCTGGCCC					
2350	2360	2370	2380	2390	2400
CAGCCCTGCG CCCTTTGCGC CCCCACGCT TGTCTGCGT GCGCTGCCCG CTCTTCATT					
2410	2420	2430	2440	2450	2460
TACCTTCTCT CCCACCAAG TTTGTACTCT TTTCTTCTC TCGGTTTTAT TTTTGTTTT					
2470	2480	2490	2500	2510	2520
TGTTTGTTTG TTTGAGACAG GCTTTCGCTC TGTCTCCAG GCTGGAGTGC AGTGGCGCGA					
2530	2540	2550	2560	2570	2580
TCTCGGCTCA CTGCAGCCTC CACCTCCCAG GTTCAAGCGA TCCGCCTGCC GAGTAGCTGG					

FIG. 1D



2590 2600 2610 2620 2630 2640
GATTACAGGC GCCCGCCACC ACGCCTGGCT AATTTTTGTG TTTTGTAGAG ATGGGGTTTC

2650 2660 2670 2680 2690 2700
GCCATGTTGG CCAGGCTGGC CTCGAACTGC TCAGCTCAAG CAATCCGCCC GCCTCGGCCT

2710 2720 2730 2740 2750 2760
CACAAAGTCC TAGAATTTTA GGCATGAGCC TCCGGGTCCG GCCTGTGCTA ATCCTTTCTG

2770 2780 2790 2800 2810 2820
TCCTTGGTTC TTTATTTCCC TTCTCTCTT TTCTTAGTCC CTTTGTCTT TCCCTCTCC

2830 2840 2850 2860 2870 2880
CGTTCAGTTG GCTGTCGTTT GAGCCTCCAC CTTTCACTC CCTCCTTTCC ACCACGATGC

2890 2900 2910 2920 2930 2940
CGAGCCCTGC CTTGGATGGG GACCATCAGC GATGACCACA ATGACCTCTC CTTACCAGG

2950 2960 2970 2980 2990 3000
CAGCTCCAGG CAGTGTTTCT GCACCGCCTT TCCCAAGGCT TGGGGGCTTT TTCTAGTGGG

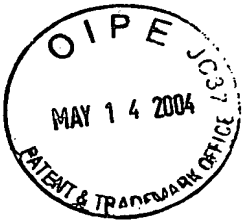
3010 3020 3030 3040 3050 3060
CTTTGAGCTG CTCAATCTGG CCTCTGCAGG GCCGGCTCCC AGCCCTTCCA ACCTCCTCAC

3070 3080 3090 3100 3110 3120
AGCCCGACCT GGGACCTAGC CAATTCCCGG AGAGTCTCTG TCCCATCGTG ACCCCCTCAC

3130 3140 3150 3160 3170 3180
AACTCTCCCA CTCACCAAAG TCTGATGACT GTGCTAGGGG GTGCTTATAT AGAGTACTGA

3190 3200 3210 3220 3230 3240
GTGTTACAAA AGCAGAAGTC TGGATGAGAA CCAATTTGTG ATATTAAGCA GGTGGGGTGG

FIG.1E



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3250 3260 3270 3280 3290 3300
GGGTGGGGAG TGTACCTAGG TTCATTTTCC GCCCTGCTTT TCCCCTTTCC AGTGTGTGCA

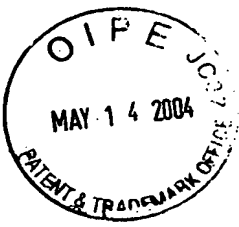
3310 3320 3330 3340 3350 3360
CTTAACCAGT CCCTGGGCCC TGTCCCCAT CCCCCTCCAA GGCATGGATT GGGTGGGCTT

3370 3380 3390 3400 3410 3420
GTGTGTCTTG GGGCAGGTGG CCCTTTCTAA ACTCTCTGCC TTTGCTCACC CACAGGACAC

3430 3440 3450 3460 3470 3480
ATAGTATGAC CATTAGGTGT TTCGTCTCCC ACCCATTTTC TATGGAAAAC CAAGGGGATC

3490 3500 3510 3520 3530 3540
GGGCCATGAT AGCCACTGGC AGCTT. (SEQ ID NO:1)

FIG. 1F



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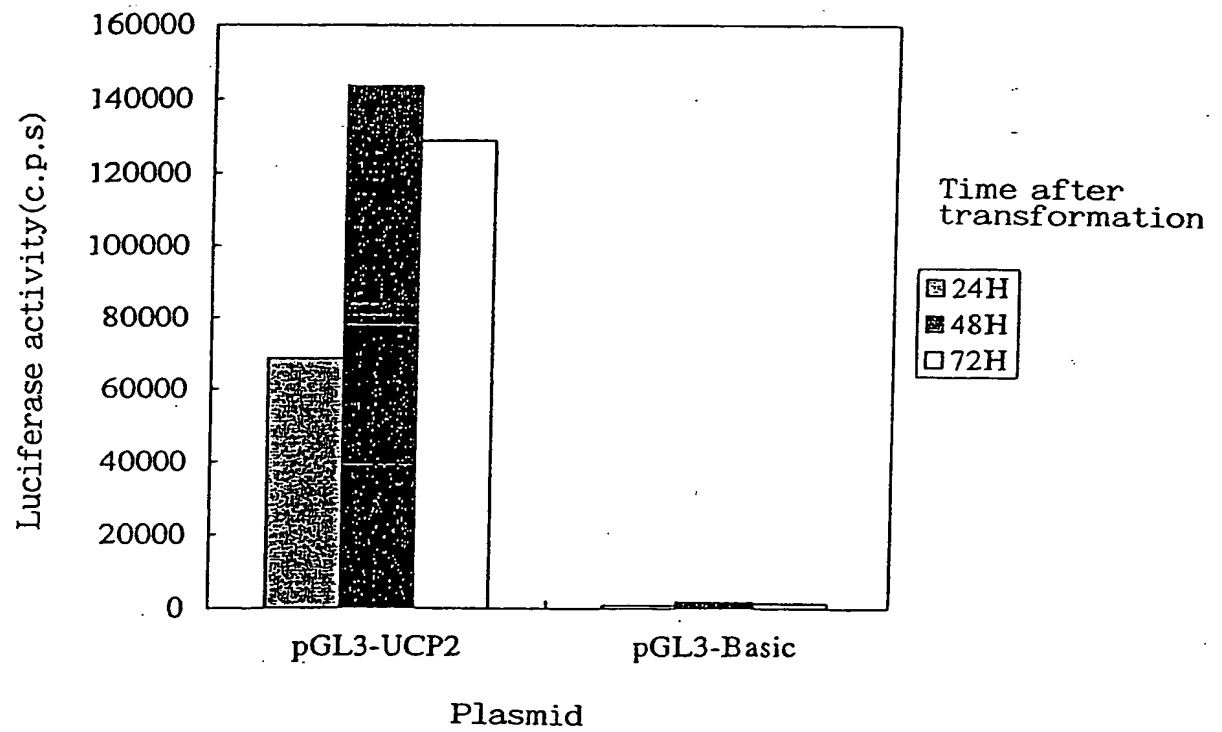
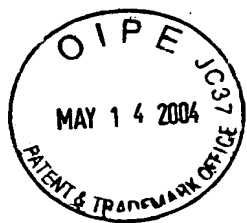


FIG. 2



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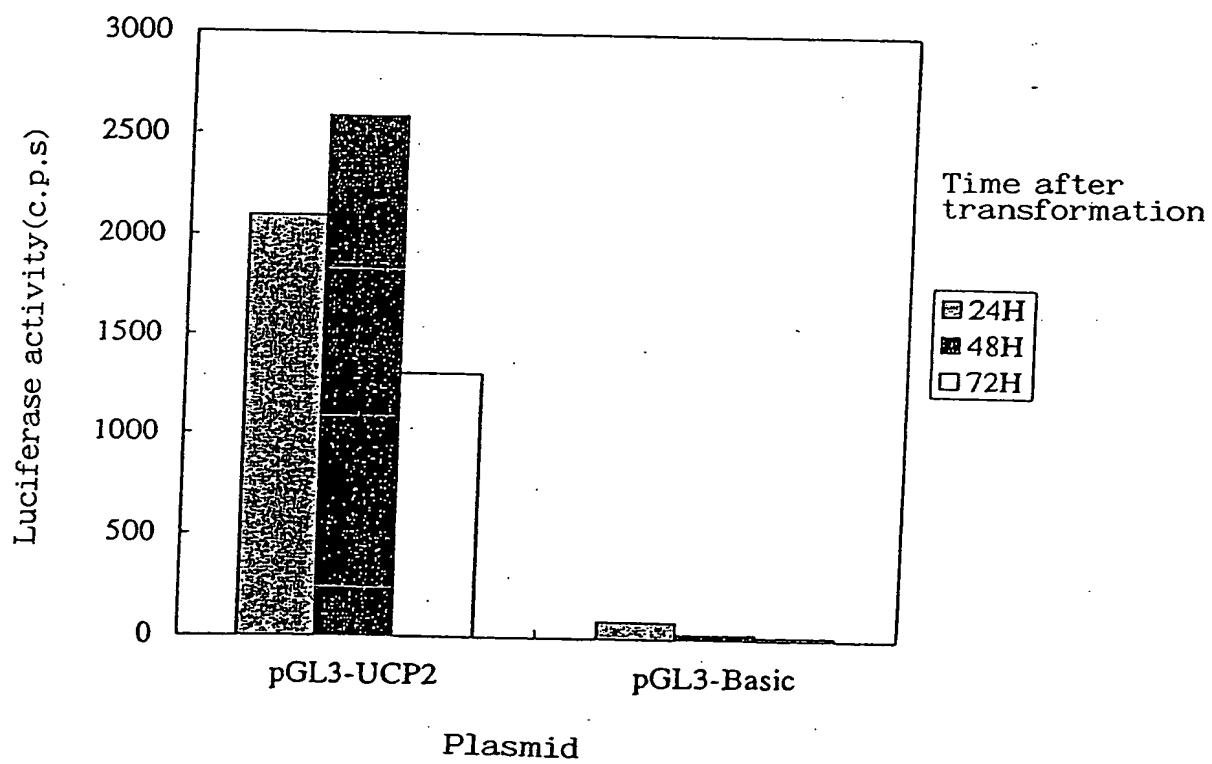


FIG. 3

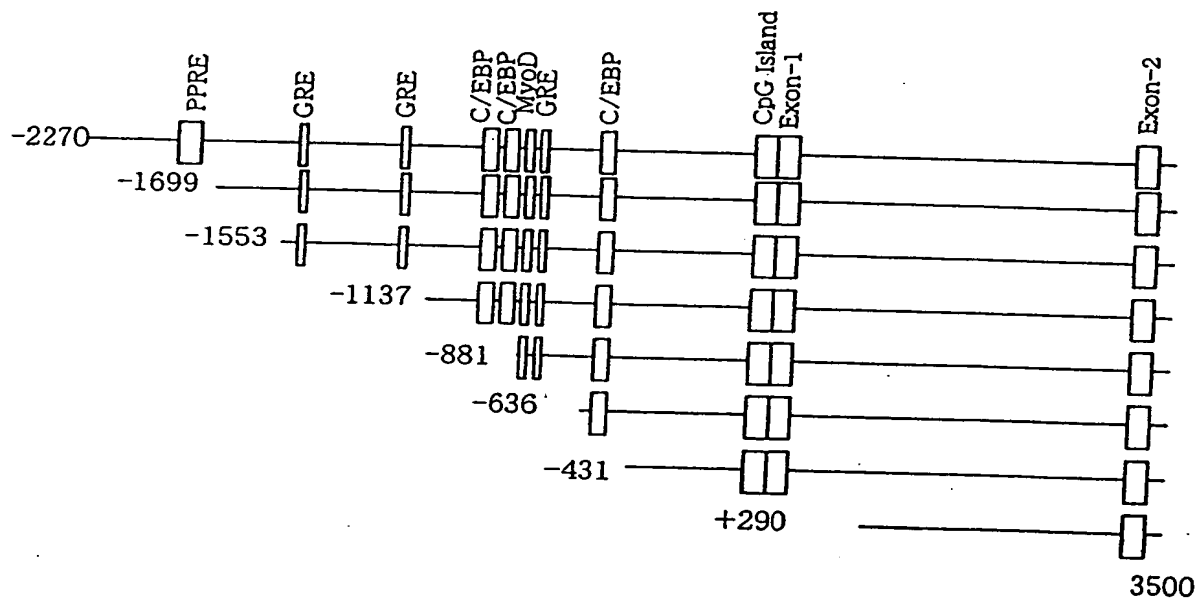
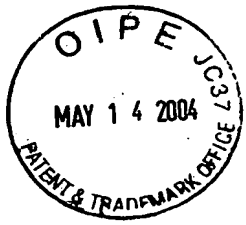


FIG. 4



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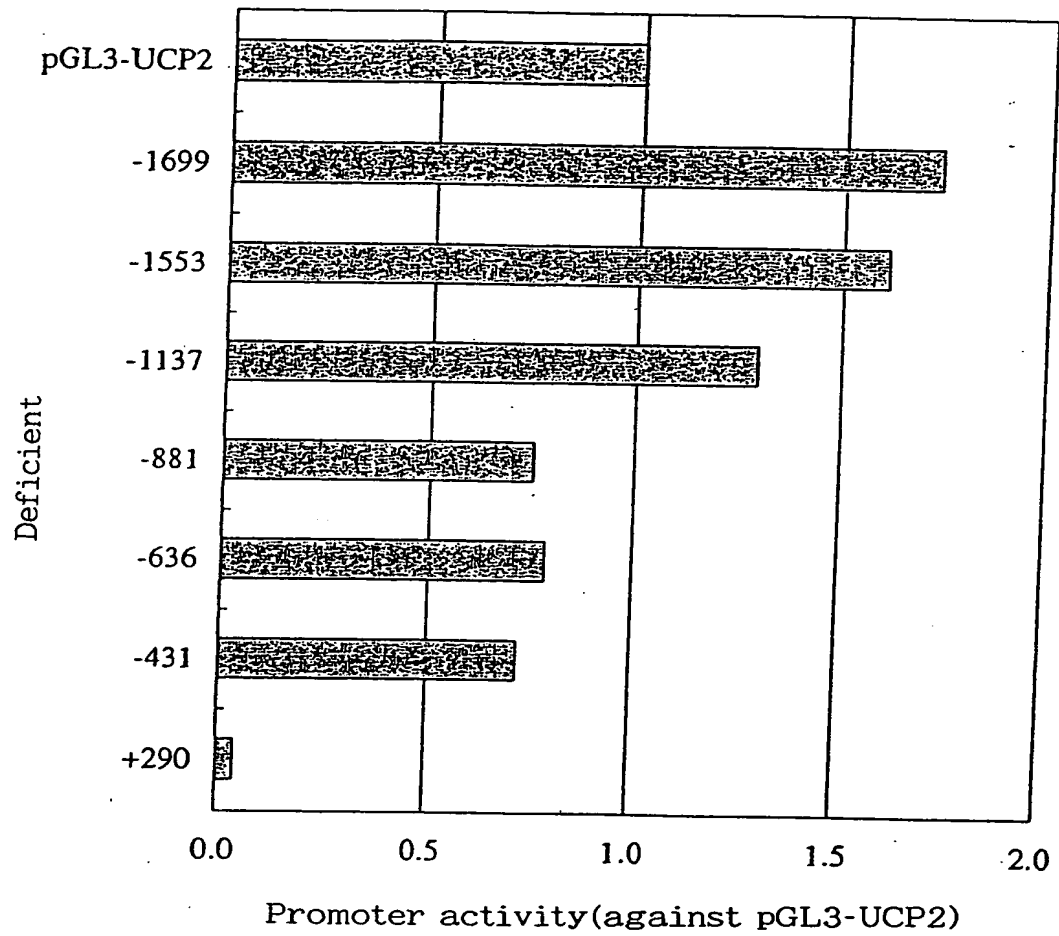


FIG. 5